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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/828,270

DATE: 03/29/2002

TIME: 13:54:52

Input Set : A:\-58-1.app

Output Set: N:\CRF3\03292002\I828270.raw

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3 <110> APPLICANT: Powers, Jay P.
4     Jaen, Juan C.
5     Piper, Derek E.
6     Walker, Nigel P.C.
7     Li, Yang
8     Tularik Inc.
10 <120> TITLE OF INVENTION: NS5B HCV Polymerase Inhibitors
12 <130> FILE REFERENCE: 018781-005810US
14 <140> CURRENT APPLICATION NUMBER: US 09/828,270
15 <141> CURRENT FILING DATE: 2001-04-05
17 <150> PRIOR APPLICATION NUMBER: US 60/194,912
18 <151> PRIOR FILING DATE: 2000-04-05
20 <160> NUMBER OF SEQ ID NOS: 1
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 591
26 <212> TYPE: PRT
27 <213> ORGANISM: Hepatitis C virus
29 <220> FEATURE:
30 <223> OTHER INFORMATION: HCV NS5B RNA-dependent RNA polymerase (RdRp)
31     (EC 2.7.7.48)
33 <400> SEQUENCE: 1
34 Ser Met Ser Tyr Thr Trp Thr Gly Ala Leu Ile Thr Pro Cys Ala Ala
35 1          5          10          15
36 Glu Glu Ser Lys Leu Pro Ile Asn Pro Leu Ser Asn Ser Leu Leu Arg
37          20          25          30
38 His His Ser Met Val Tyr Ser Thr Ser Arg Ser Ala Ser Leu Arg
39          35          40          45
40 Gln Lys Lys Val Thr Phe Asp Arg Leu Gln Val Leu Asp Asp His Tyr
41          50          55          60
42 Arg Asp Val Leu Lys Glu Met Lys Ala Lys Ala Ser Thr Val Lys Ala
43 65          70          75          80
44 Arg Leu Leu Ser Ile Glu Glu Ala Cys Lys Leu Thr Pro Pro His Ser
45          85          90          95
46 Ala Lys Ser Lys Phe Gly Tyr Gly Ala Lys Asp Val Arg Ser Leu Ser
47          100         105         110
48 Ser Arg Ala Val Asn His Ile Arg Ser Val Trp Glu Asp Leu Leu Glu
49          115         120         125
50 Asp Thr Glu Thr Pro Ile Asp Thr Thr Ile Met Ala Lys Asn Glu Val
51          130         135         140
52 Phe Cys Val Gln Pro Glu Lys Gly Gly Arg Lys Pro Ala Arg Leu Ile
53 145         150         155         160
54 Val Phe Pro Asp Leu Gly Val Arg Val Cys Glu Lys Met Ala Leu Tyr

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55                               165                               170                               175
56 Asp Val Val Ser Thr Leu Pro Gln Ala Val Met Gly Pro Ser Tyr Gly
57                               180                               185                               190
58 Phe Gln Tyr Ser Pro Gly Gln Arg Val Glu Phe Leu Val Asn Thr Trp
59                               195                               200                               205
60 Lys Ser Lys Lys Cys Pro Met Gly Phe Ser Tyr Asp Thr Arg Cys Phe
61                               210                               215                               220
62 Asp Ser Thr Val Thr Glu Asn Asp Ile Arg Thr Glu Glu Ser Ile Tyr
63 225                               230                               235                               240
64 Gln Cys Cys Asp Leu Ala Pro Glu Ala Arg Gln Ala Ile Arg Ser Leu
65                               245                               250                               255
66 Thr Glu Arg Leu Tyr Val Gly Gly Pro Leu Thr Asn Ser Lys Gly Gln
67                               260                               265                               270
68 Asn Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr Ser
69                               275                               280                               285
70 Cys Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Cys Arg
71                               290                               295                               300
72 Ala Ala Lys Leu Gln Asp Cys Thr Met Leu Val Asn Gly Asp Asp Leu
73 305                               310                               315                               320
74 Val Val Ile Cys Glu Ser Ala Gly Thr Gln Glu Asp Ala Ala Ala Leu
75                               325                               330                               335
76 Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser Ala Pro Pro Gly Asp
77                               340                               345                               350
78 Pro Pro Gln Pro Glu Tyr Asp Leu Glu Leu Ile Thr Ser Cys Ser Ser
79                               355                               360                               365
80 Asn Val Ser Val Ala His Asp Ala Ser Gly Lys Arg Val Tyr Tyr Leu
81                               370                               375                               380
82 Thr Arg Asp Pro Thr Thr Pro Leu Ala Arg Ala Ala Trp Glu Thr Val
83 385                               390                               395                               400
84 Arg His Thr Pro Val Asn Ser Trp Leu Gly Asn Ile Ile Met Tyr Ala
85                               405                               410                               415
86 Pro Thr Leu Trp Ala Arg Met Ile Leu Met Thr His Phe Phe Ser Ile
87                               420                               425                               430
88 Leu Leu Ala Gln Glu Gln Leu Glu Lys Ala Leu Asp Cys Gln Ile Tyr
89                               435                               440                               445
90 Gly Ala Cys Tyr Ser Ile Glu Pro Leu Asp Leu Pro Gln Ile Ile Glu
91                               450                               455                               460
92 Arg Leu His Gly Leu Ser Ala Phe Ser Leu His Ser Tyr Ser Pro Gly
93 465                               470                               475                               480
94 Glu Ile Asn Arg Val Ala Ser Cys Leu Arg Lys Leu Gly Val Pro Pro
95                               485                               490                               495
96 Leu Arg Val Trp Arg His Arg Ala Arg Ser Val Arg Ala Lys Leu Leu
97                               500                               505                               510
98 Ser Gln Gly Gly Arg Ala Ala Thr Cys Gly Lys Tyr Leu Phe Asn Trp
99                               515                               520                               525
100 Ala Val Lys Thr Lys Leu Lys Leu Thr Pro Ile Pro Ala Ala Ser Gln
101                               530                               535                               540
102 Leu Asp Leu Ser Gly Trp Phe Val Ala Gly Tyr Asn Gly Gly Asp Ile
103 545                               550                               555                               560

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104	Tyr	His	Ser	Leu	Ser	Arg	Ala	Arg	Pro	Arg	Trp	Phe	Met	Leu	Cys	Leu
105					565					570					575	
106	Leu	Leu	Leu	Ser	Val	Gly	Val	Gly	Ile	Tyr	Leu	Leu	Pro	Asn	Arg	
107					580					585					590	

VERIFICATION SUMMARY

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